

**Exhibit A**

**Clean Version of The Pending Claims in U.S. Patent Application Ser. No. 09/975,308**

1. (Amended) An isolated expression vector comprising the nucleotide sequence of SEQ ID NO:8.
2. (Amended) An isolated expression vector comprising a nucleic acid sequence encoding the amino acid sequence of SEQ ID NO:9.
3. (New) A host cell comprising the recombinant expression vector of claim 1 or 2.

BD144530 ACCESSION:BD144530 NID: gi 27850288 dbj BD144530.1 Novel  
G-protein coupled receptors  
Length = 924

Score = 617 bits (1574), Expect = e-174  
Identities = 305/307 (99%), Positives = 307/307 (99%)  
Frame = +1

Query: 1 MNHSVVTEFIILGLTKKPELQGIIIFLFFLIVYLVAFLGNMLIIIAKIYSNTLHTPMYVFL 60  
MNHSVVTEFIILGLTKKPELQGIIIFLFFLIVYLVAFLGNMLIIIAKIY+NTLHTPMYVFL  
Sbjct: 1 MNHSVVTEFIILGLTKKPELQGIIIFLFFLIVYLVAFLGNMLIIIAKIYNNLHTPMYVFL 180

Query: 61 LTLAVVDIICTTSIIPKMLGTMLTSENTISYAGCMSQLFLFTWSLGAEMVLFTTMAYDRY 120  
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Sbjct: 181 LTLAVVDIICTTSIIPKMLGTMLTSENTISYAGCMSQLFLFTWSLGAEMVLFTTMAYDRY 360

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VAICFPLHYST+MNHHMCVALLSMVMAIAVTNSWVHTALIMRLTFCGPNTIDHFFCEIPP  
Sbjct: 361 VAICFPLHYSTVMNHHMCVALLSMVMAIAVTNSWVHTALIMRLTFCGPNTIDHFFCEIPP 540

Query: 181 LLALSCSPVRINEVMVYVADITLAIGDFILTCISYGFIIIVAILRIRTVEGKRKAFSTCSS 240  
LLALSCSPVRINEVMVYVADITLAIGDFILTCISYGFIIIVAILRIRTVEGKRKAFSTCSS  
Sbjct: 541 LLALSCSPVRINEVMVYVADITLAIGDFILTCISYGFIIIVAILRIRTVEGKRKAFSTCSS 720

Query: 241 HLTVVTLYYSPVIYTYIRPASSYTFERDKVVAALYTLVPTLNPMVYSFQNREMQAGIRK 300  
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Sbjct: 721 HLTVVTLYYSPVIYTYIRPASSYTFERDKVVAALYTLVPTLNPMVYSFQNREMQAGIRK 900

Query: 301 VFAFLKH 307  
VFAFLKH  
Sbjct: 901 VFAFLKH 921



**Nucleotide**

PubMed

Nucleotide

Protein

Genome

### Structure

PMc

### Taxonomy

OMIM

Boo

Search  for

### Limits

[Preview/Index](#)

## History

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## Details

## Display

default

Show:

20

Send to

File

## Get Subsequence

□ 1: BD144530. Novel G-protein c...[gi:27850288]

## Links

```

LOCUS       BD144530                      924 bp    DNA        linear        PAT 17-JAN-2003
DEFINITION  Novel G-protein coupled receptors.
ACCESSION   BD144530
VERSION     BD144530.1  GI:27850288
KEYWORDS    JP 2002112793-A/255.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
              Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 924)
  AUTHORS   Haga,T., Takeda,S. and Miyake,N.
  TITLE     Novel G-protein coupled receptors
  JOURNAL   Patent: JP 2002112793-A 255 16-APR-2002;
            JAPAN SCIENCE AND TECHNOLOGY CORP
COMMENT     OS   Homo sapiens (human)
            PN   JP 2002112793-A/255
            PD   16-APR-2002
            PF   09-FEB-2001 JP 2001034434
            PI   TATSUYA HAGA, SHIGEKI TAKEDA, NARIKI MIYAKE
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              G01N33/566//
            PC   C12P21/08,C12N15/00,A61K37/02,C12N5/00
            CC   Novel G-protein coupled receptors
            FH   Key                      Location/Qualifiers
            FT   CDS                      (1)..(924).

FEATURES             Location/Qualifiers
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901 gtgtttgcat ttctgaaaca ctag
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//

Revised: July 5, 2002.

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Feb 19 2003 14:49:48

Query= SEQ ID NO:8  
 (924 letters)

Sequences producing significant alignments:		Score	E
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Strand = Plus / Minus			
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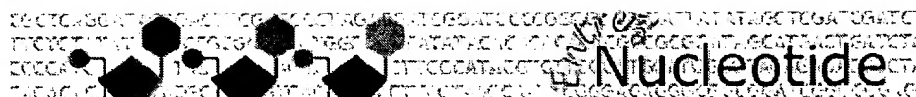
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PubMed

Nucleotide

Protein

Genome

Structure

PMC

Taxonomy

OMIM

Boo

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Show:

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☐ 1: AC091612. Homo sapiens chro...[gi:18497169]

Links

LOCUS AC091612 180657 bp DNA linear HTG 05-FEB-2002  
 DEFINITION Homo sapiens chromosome 1 clone RP11-656022, WORKING DRAFT  
 SEQUENCE, 1 unordered piece.  
 ACCESSION AC091612 AL390860  
 VERSION AC091612.4 GI:18497169  
 KEYWORDS HTG; HTGS\_PHASE1; HTGS\_DRAFT; HTGS\_FULLTOP.  
 SOURCE Homo sapiens (human)  
 ORGANISM Homo sapiens  
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
 REFERENCE 1 (bases 1 to 180657)  
 AUTHORS Kaul,R.K., Olson,M.V., Raymond,C. and Haugen,E.D.  
 TITLE Direct Submission  
 JOURNAL Unpublished  
 REFERENCE 2 (bases 1 to 180657)  
 AUTHORS Kaul,R.K., Olson,M.V., Raymond,C., Clendenning,J., Ivey,R.G. and  
 Haugen,E.D.  
 TITLE Direct Submission  
 JOURNAL Submitted (09-MAY-2001) Genome Center, University of Washington,  
 Box 352145, Seattle, WA 98195, USA  
 COMMENT On Feb 5, 2002 this sequence version replaced gi:15487406.  
 ----- Genome Center  
 Center: University of Washington Genome Center  
 Center Code: UWGC  
 Web site: <http://www.genome.washington.edu>  
 Contact: uwgchtgs@u.washington.edu  
 Drafting Center: SC  
 ----- Project Information  
 Center project name: chr-1  
 Center clone name: RP11-656022 (sc0182)  
 ----- Summary Statistics  
 Sequencing vector: plasmid; L08752; 100% of reads  
 Chemistry: Dye-terminator Big Dye; 100% of reads  
 Assembly program: Phrap; version 0.990319  
 Consensus quality: 180536 bases at least Q40  
 Consensus quality: 180650 bases at least Q30  
 Consensus quality: 180657 bases at least Q20  
 Insert size: 194815; 11.0% error; agarose-fp  
 Insert size: 180657; sum-of-contigs  
 Quality coverage: 8.4x in Q20 bases; agarose-fp  
 Quality coverage: 9.0x in Q20 bases; sum-of-contigs  
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 \* NOTE: This is a 'working draft' sequence. It currently  
 \* consists of 1 contigs. The true order of the pieces  
 \* is not known and their order in this sequence record is  
 \* arbitrary. Gaps between the contigs are represented as  
 \* runs of N, but the exact sizes of the gaps are unknown.